

### **BLAST 2 SEQUENCES**

This tool produces the alignment of two given sequences using <u>BLAST</u> engine for local alignment.

The stand-alone executable for blasting two sequences (bl2seq) can be retrieved from <u>NCBI ftp site</u>

Reference: Tatiana A. Tatusova, Thomas L. Madden (1999), "Blast 2 sequences - a new tool for comparing protein and nucleotide sequences", FEMS Microbiol Lett. 174:247-250

Program blastp Matrix BLOSUM62	
Parameters used in <u>BLASTN</u> program only:  Reward f r a match: Penalty for a mismatch:	
Use Mega BLAST Strand option Not Applicable	
Open gap 11 and extension gap 1 penalties gap x_dropoff 50 expect 10.0 word size 3 Filter Align	
Sequence 1 Enter accession or GI SEQ 4 or download from file	
or sequence in FASTA format from: 0 to: 0	
qsegpavvnıqaapaprtqngsgnaetasapıaqsapryerikrivpnmpelpqeeadagg lnfgsgfiiskngyiltnthvvagmgsikvllndkreytakligsdvqsdvallkidatee lpvvkignpknlkpgewvaaigapfgfdnsvtagivsakgrslpnesytpfiqtdvainpg nsggplfnlkgqvvginsqiysrsggfmgisfaipidvamnvaeqlkntgkvqrgqlgvii qevsyglaqsfgldkasgaliakilpgspaeraglqagdivlsldggeirssgdlpvmvga itpgkevslgvwrkgeeitikaklgnaaehtgassktdeapyteqqsgtfsvesagitlqt htdssgkhlvvvrvsdaaeraglrhgdeilavrasprq	
Sequence 2 Enter accession or GI Gilbert4 or download from file	· · · · · · · · · · · · · · · · · · ·
or sequence in FASTA format from: 0 to: 0  mgikkvcitviciivicigirytrarvhqqernavsiikakiineegkpvniitcytiiqm kvaerimaqhpqerfyvvlmsenrnekydyyfnqikdkaerayffylpyglnksfnfiptm aelkvksmllpkvkriylaslekvsiaaflstypdaeiktfddgtnnliressylggefav ngaikrnfarmmvgdwsiaktrnasdehytifkglknimddgrrkmtylplfdaselkagd etggtvrillgspdkemkeisekaaknfniqyvaphprqtyglsgvtalnspyviedyilr eikknphtryeiytffsgaaltmkdfpnvhvyalkpaslpedywlkpvyalfrqadipilt fddkn	
Align Clear Input	

Comments and suggestions to <u>blast-help@ncbi.nlm.nih.gov</u>



# Blast 2 Sequences results

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.4 [Aug-26-2002]

Matrix BLOSUM62 ap gap open: 11 gap extension: 1

x\_dropoff: 50 expect: 10.0 wordsize: 3 Filter ✓ Align

Sequence 1 |cl||seq\_1 | Length 465 | O9/1839, O90 | SEQ 10 NOTY

Sequence 2 |cl||seq\_2 | Length 371 | Cot O96, 529 | SEQ 10 NOT 2

N significant similarity was found

### **BLAST 2 SEQUENCES**

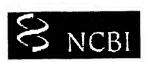
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ogram blastp Matrix BLOSUM62
eward f r a match:  Penalty for a mismatch:
Use Mega BLAST Strand option Not Applicable
pen gap 11 and extension gap 1 penalties  ap x_dropoff 50 expect 10.0 word size 3 Filter Align
equence 1 Enter accession or GI SEQ 4 or download from file
sequence in FASTA format from: 0 to: 0
segpavvnıqaapaprtqngsgnaetdsdpiadsdpiyerikrivpnmpelpqeeaddgg infgsgfiiskngyiltnthvvagmgsikvllndkreytakligsdvqsdvallkidatee pvvkignpknlkpgewvaaigapfgfdnsvtagivsakgrslpnesytpfiqtdvainpg isggplfnlkgqvvginsqiysrsggfmgisfaipidvamnvaeqlkntgkvqrgqlgvii gevsyglaqsfgldkasgaliakilpgspaeraglqagdivlsldggeirssgdlpvmvga itpgkevslgvwrkgeeitikaklgnaaehtgassktdeapyteqqsgtfsvesagitlqt itdssgkhlvvvrvsdaaeraglrhgdeilavrasprq
equence 2 Enter accession or GI Gilbert2 or download from file
r sequence in FASTA format from: 0 to: 0
ngikkacitviciivicigiiytiqrvnngernavsiikokiineegepvniiicytiiqm kvaerimaqhpgerfyvvlmsenrnekydyyfkqikdkaerayffhlpyglnksfnfiptm nelkvksmllpkvkriylaslekvsiaaflstypdaeiktfddgtgnliqsssylgdefsv ngtikrnfarmmigdwsiaktrnasdehytifkglknimddgrrkmtylplfdaselkagd etggtvrillgspdkemkeisekaaknfniqyvaphprqtyglsgvttlnspyviedyilr eikknphtryeiytffsgaaltmkdfpnvhvyalkpaslpedywlkpvyalftqsgipilt fddkn
Align Clear Input

Comments and suggestions to <u>blast-help@ncbi.nlm.nih.gov</u>



# Blast 2 Sequ nc s results

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.4 [Aug-26-2002]

Matrix BLOSUM62 gap open: 11 gap extension: 1 x\_dropoff: 50 expect: 10.0 wordsize: 3 Filter Malign

Sequence 1 Icl|seq\_1 Length 465 09/3PP, 090 3EG D VOY

No significant similarity was found

Sequence 2 |cl|seq\_2 | Length 371 | 6 096, 529 | Sta 10 NO: 4



Express Mail No.: EL 500 578 282 US

#### IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of Jackson and Horris

RECEIVED

DEC 2 7 2002

Serial No.: 09/388,090

Art Unit: 1645

Filed: August 31, 1999

Examiner: S. Devi

**TECH CENTER 1600/2900** 

For: NEISSERIA SPP.

Anomey Ducket No.: 7969-082-999

POLYPEPTIDE, NUCLEIC ACID

SEQUENCE AND USES

THEREOF

### Declaration of Br. W. James Jackson Under C.F.R. 8 1.132

Assistant Commissioner for Patents Washington, D.C. 20231

SIR:

#### L.Dr. W. James Jackson, declare and state:

- I am a co-inventor of the above-identified application which I have read and 1. understand.
- The following experiments were conducted under my supervision and control. The 2. results obtained are shown in Table 2 below. As detailed below, the results obtained olearly demonstrate that the isolated Neisseria polypeptide designated "NGSP" polypeptide taught in the application is useful to induce an immune response against infection by Neisseria gonorrhea.
- Methods: The murine model of vaginal colonization was used to evaluate the 3. ability of the isolated NGSP polypoptide of the invention, as described in Jerse, 1999, Infec. Immun. 67(11): 5699-5708, a copy of which is attached as Exhibit A, as modified as detailed below.
- 3a. 'The NGSP polypeptide (containing a His tag) was recombinately expressed in E. coli JM109 containing plasmid pTLZ-NgHtr A#2 (ATCC number PTA-470) (688

Sections 8.3 and 8.4 of the application) and isolated using affinity chromotograph as described in the application in Section 5.3.

- 3b. Balb/o female mice, six to ten weeks of age, were immunized three times at two week intervals with isolated recombinant NGSP polypeptide. Animals were immunized, parenterally (s.c.) according to the groupings defined in Table 1 below.
- 3c. Via parenteral immunization, the NGSP polypoptide was coadministered with Freund's complete adjuvant (CFA) at the first immunization. An equal volume of CFA was mixed with the NGSP polypoptide antigen solution to form an emulsion. At the second and third immunizations, Freund's incomplete adjuvant (IFA) was co-administered with the NGSP polypoptide. An equal volume of IFA was mixed with the NGSP polypoptide antigen solution to form an emulsion. Animals immunized three times with PBS alone served as negative controls, while animals immunized with formalia treated Netweria governers whole cell antigen served as positive controls.

<u>Table 1</u>

Group Number	Autigen Amount per Dose	Adjuvant
1	NGSP polypeptide (50)	CFA/IFA
2 positive control	Inactivated Neusseria gonnorrhea whole cell antigen	CFA/IFA
3 negative control	PBS Only	None

- 1 Antigen dose is given in micrograms (µg).
- 3d. Retrooribital bleeds were taken prior to the first immunization and again approximately ten days following the third and final dose. Approximately ten days after the last immunization, a slow release pellet of  $\beta$ -estradiol (5.0mg released over 21 days; Innovative Research Inc.) was implanted under the skin of each animal to stabilize and synchronize the uterine cycle prior to and during the challenge phase of the experiment. Animals were challenged three to five days later by instilling into the vaginal canal a 20  $\mu$ l volume of PBS containing -1.0 X  $10^6$  cfn of the N. genorrhea strain MS11A.

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- 3c. For challenge, N. gonnorhea strain MS11A was grown from frozen stock cultures on Thayer Martin chocolate agar plates, collected by swabbing the sterile applicators and suspended in PBS. The N. gonnorhea strain MS11A suspension was diluted with PBS to ~1.0 X 10<sup>8</sup> cfu/ml using a previously generated O.D. so vs cfu standard curve.
- 3f. After challenge (48hrs post challenge) the level of N. gonnorhea vaginal colonization was assessed by swabbing the vaginal canal with pediatric nasopharengyl Dacron/Polyester swabs. Swabs were inserted gently into the canal until resistance was encountered then gently rotated (10 complete 360° turns) to collect infectious microorganisms. Swabs were removed, moistened by brief immersion into sterile PBS and used to streak plate Thayer Martin chocolate agar plates containing antibiotics (vancomycin, collistin, nystatin, timethoprim). A standard quantitative dilution streak plating mathod was used to enumerate colonies. Plates were incubated at 37°C under oxygen depleted or microaerophilic conditions for 24 to 72 hours prior to counting.

Protective efficacy is expressed as percent of animals protected in a vaccination group.

Results: Results are presented in Table 2 below.

Table 2

Group Number	Ratio of Intected to Total Animals	Percent Protected
1	4/10	60%
2	3/10	70%
3	6/8	25%

5. As illustrated in Table 2, fewer mice immunized with isolated recombinant NGSP polypeptide were found to be infected with *N. gonnorhea* after challenge compared to the unimmunized and unprotected negative controls. The level of protection conferred by NGSP polypeptide of the invention is equivalent to that obtained with the positive control, i.e., inactivated *N. gonnorrhea* whole cell antigen. These results clearly demonstrate that isolated NGSP polypeptide of the present application can confer protection against vaginal

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infection by N. gonnorhea. These results demonstrate that the isolated NGSP polypeptide is useful to induce a protective immune response against Neisseria generrhea in a relevant animal model and thus is useful for a protective human vaccine.

I declare further that all statements made in this Declaration of my own knowledge are true and that all statements made on information and helief are helieved to be true and further that these statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the specification or any patent issuing thereon.

Dated

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